
Sequence Listing could not be accepted.

If you need help call the Patent Electronic Business Center at (866)

217-9197 (toll free).

Reviewer: Anne Corrigan

Timestamp: Fri Sep 07 16:38:33 EDT 2007

Reviewer Comments:

SEQUENCE LISTING

<110> MADDON, Paul J.

DONOVAN, Gerald P.

OLSON, William C.

SCHÜLKE, Norbert

GARDNER, Jason

MA, Dangshe

Please remove all above foreign accent marks—they are non-ASCII characters and cannot be processed.

<210> 14

<211> 481

<212> DNA

<213> Artificial Sequence

<220>

<223> Includes BamHI/Bg1II cloning junction, signal peptide, V region,
portion of C region and 3'XbaI/NheI (heavy) or NheI (light) cloning
junction

The above <223> response exceeds 72 characters per line; per 1.823 of the Sequence Rules, a maximum of 72 characters per line, including white spaces. This line-length problem appears in subsequent sequences, too.

Validated By CRFValidator v 1.0.3

Application No: 10695667 Version No: 2.0

Input Set:

Output Set:

Started: 2007-08-27 11:46:11.987

Finished: 2007-08-27 11:46:18.117

Elapsed: 0 hr(s) 0 min(s) 6 sec(s) 130 ms

Total Warnings: 26

Total Errors: 0

No. of SeqIDs Defined: 37

Actual SeqID Count: 37

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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(3)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(4)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(5)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(6)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(7)
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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(9)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(10)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(11)
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W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(24)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(26)
W	213	Artificial	or	Unknown	found	in	<213>	in	SEQ	ID	(28)

Input Set:

Output Set:

Started: 2007-08-27 11:46:11.987

Finished: 2007-08-27 11:46:18.117

Elapsed: 0 hr(s) 0 min(s) 6 sec(s) 130 ms

Total Warnings: 26

Total Errors: 0

No. of SeqIDs Defined: 37

Actual SeqID Count: 37

Error code Error Description

This error has occured more than 20 times, will not be displayed

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<213> Homo sapiens
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     35
             40
                                45
Ala Thr Asn Ile Thr Pro Lys His Asn Met Lys Ala Phe Leu Asp Glu
  50 55 60
Leu Lys Ala Glu Asn Ile Lys Lys Phe Leu Tyr Asn Phe Thr Gln Ile
        70
                  75
Pro His Leu Ala Gly Thr Glu Gln Asn Phe Gln Leu Ala Lys Gln Ile
                  90 95
          85
Gln Ser Gln Trp Lys Glu Phe Gly Leu Asp Ser Val Glu Leu Ala His
      100 105
Tyr Asp Val Leu Leu Ser Tyr Pro Asn Lys Thr His Pro Asn Tyr Ile
    115
             120
                           125
Ser Ile Ile Asn Glu Asp Gly Asn Glu Ile Phe Asn Thr Ser Leu Phe
  130 135 140
Glu Pro Pro Pro Gly Tyr Glu Asn Val Ser Asp Ile Val Pro Pro
        150 155
145
                                   160
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Ser Tyr Pro Asp Gly Trp Asn Leu Pro Gly Gly Gly Val Gln Arg Gly 245 250 255

Asn Ile Leu Asn Leu Asn Gly Ala Gly Asp Pro Leu Thr Pro Gly Tyr 260 265 270

Pro Ala Asn Glu Tyr Ala Tyr Arg Arg Gly Ile Ala Glu Ala Val Gly 275 280 285

Leu Pro Ser Ile Pro Val His Pro Ile Gly Tyr Tyr Asp Ala Gln Lys 290 295 300

Leu Leu Glu Lys Met Gly Gly Ser Ala Pro Pro Asp Ser Ser Trp Arg 305 310 315 320

Gly Ser Leu Lys Val Pro Tyr Asn Val Gly Pro Gly Phe Thr Gly Asn 325 330 335

Phe Ser Thr Gln Lys Val Lys Met His Ile His Ser Thr Asn Glu Val 340 345 350

Thr Arg Ile Tyr Asn Val Ile Gly Thr Leu Arg Gly Ala Val Glu Pro 355 360 365

Asp Arg Tyr Val Ile Leu Gly Gly His Arg Asp Ser Trp Val Phe Gly 370 375 380

Gly Ile Asp Pro Gln Ser Gly Ala Ala Val Val His Glu Ile Val Arg 385 390 395 400

Ser Phe Gly Thr Leu Lys Lys Glu Gly Trp Arg Pro Arg Arg Thr Ile
405 410 415

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Asp	Lys 610	Ile	Tyr	Ser	Ile	Ser 615	Met	Lys	His	Pro	Gln 620	Glu	Met	Lys	Thr
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480

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<213> Artificial Sequence

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